# This listing of claims will replace all prior versions and listings of claims in the application: Listing of Claims

- 1. (Original) A composition comprising a nucleic acid wherein the nucleic acid comprises a sequence encoding a HEX- $\alpha$  and a sequence encoding a HEX- $\beta$ .
- 2. (Original) The composition of claim 1, wherein the sequence encoding the HEX- $\beta$  is orientated 5' to the sequence encoding HEX- $\alpha$ .
  - 3. (Original) The composition of claim 1, further comprising a promoter.
- 4. (Original) The composition of claim 1, further comprising an integrated ribosomal entry site (IRES).
- 5. (Original) The composition of claim 4, wherein the sequence encoding the HEX- $\beta$  is orientated 5' to the IRES sequence and the IRES sequence is located 5' to the sequence encoding HEX- $\alpha$ .
  - 6. (Original) The composition of claim 4, further comprising a promoter.
- 7. (Original) The composition of claim 6, wherein the promoter is located 5' to the sequence encoding the HEX- $\beta$  and the sequence encoding the HEX- $\beta$  is orientated 5' to the IRES sequence and the IRES sequence is located 5' to the sequence encoding HEX- $\alpha$ .
- 8. (Original) The composition of claim 6, wherein the parts are oriented 5'-promoter- HEX-β encoding sequence-IRES- HEX-α encoding sequence-3'.
- 9. (Original) The composition of claim 6, wherein the parts are oriented 5'-promoter- HEX-α encoding sequence -IRES- HEX-β encoding sequence -3'.
- 10. (Original) The composition of claim 6, wherein the nucleic acid comprises a second IRES sequence.

- 11. (Original) The composition of claim 10, wherein the second IRES sequence is located 3' to the other parts.
- 12. (Original) The composition of claim 6, wherein the HEX- $\beta$  has at least 70%, 75%, 80%, 85%, 90%, or 95% identity to the sequence set forth in SEQ ID NO:3 and the HEX- $\alpha$  has at least 70%, 75%, 80%, 85%, 90%, or 95% identity to the sequence set forth in SEQ ID NO:1.
- 13. (Original) The composition of claim 12, wherein any change from SEQ ID NO:3 or SEQ ID NO:1 is a conservative change.
- 14. (Original) The composition of claim 13 wherein the HEX- $\beta$  has the sequence set forth in SEQ ID NO:3 and the HEX- $\alpha$  has the sequence set forth in SEQ ID NO:1.
- 15. (Original) The composition of claim 6, wherein the sequence encoding HEX-β hybridizes to SEQ ID NO:2 under stringent conditions and wherein the HEX-α element hybridizes to SEQ ID NO:4 under stringent conditions.
- 16. (Original) The composition of claim 12, wherein the IRES sequence comprises a sequence having at least 70%, 75%, 80%, 85%, 90%, or 95% identity to the sequence set forth in SEQ ID NO:5.
- 17. (Original) The composition of claim 16, wherein the promoter sequence comprises a constitutive promoter.
- 18. (Original) The composition of claim 17, wherein the promoter sequence comprises a CMV promoter.
- 19. (Original) The composition of claim 18, wherein the CMV promoter comprises the sequence set forth in SEQID NO:32.

- 20. (Original) The composition of claim 16, wherein the promoter sequence comprises a beta actin promoter.
- 21. (Original) The composition of claim 20, wherein the beta actin promoter sequence comprises an avian beta actin promoter sequence.
- 22. (Original) The composition of claim 21, wherein the beta actin promoter sequence comprises a mammalian beta actin promoter sequence.
- 23. (Original) The composition of claim 21, wherein the beta actin promoter comprises the sequence set forth in SEQ ID NO:26.
- 24. (Original) The composition of claim 16, wherein the promoter sequence comprises an inducible promoter.
- 25. (Original) The composition of claim 18, wherein the promoter sequence further comprises a beta actin promoter.
- 26. (Original) The composition of claim 6, wherein the composition produces a functional HEXB product.
- 27. (Original) The composition of claim 6, wherein the composition produces a functional HEXA product.
- 28. (Original) The composition of claim 6, wherein the composition produces a functional HEXS product.
- 29. (Original) The composition of claim 26, wherein the composition is capable of cross correcting.
- 30. (Original) The composition of claim 26, wherein the function is the catabolism of GM2 gangliosides in mammalian cells.
- 31. (Original) The composition of claim 6, wherein the nucleic acid further comprises a reporter gene.

- 32. (Original) The composition of claim 31, wherein the reporter gene is a lacZ gene.
- 33. (Original) The composition of claim 31, wherein the reporter gene is flanked by recombinase sites.
- 34. (Original) The composition of claim 33, wherein the recombinase sites are for the cre recombinase.
- 35. (Original) The composition of claim 6, wherein the nucleic acid further comprises a transcription termination site.
- 36. (Original) The composition of claim 35, wherein the transcription termination site is oriented 5' to the promoter sequence.
- 37. (Original) The composition of claim 36, wherein the transcription termination site is flanked by recombinase sites.
- 38. (Original) The composition of claim 37, wherein the recombinase sites are for the cre recombinase.
  - 39. (Original) The composition of claim 6, further comprising a vector.
- 40. (Original) The composition of claim 39, wherein the vector comprises a lentiviral vector.
- 41. (Original) The composition of claim 40, wherein the lentiviral vector comprises a feline immunodeficiency virus.
- 42. (Original) The composition of clam 40, wherein the lentiviral vector comprises a human immunodeficiency virus.
- 43. (Original) The composition of claim 39, wherein the vector can be stably integrated for at least three months.
- 44. (Withdrawn) A composition comprising a cell wherein the cell comprises the nucleic acid of claim 6.

- 45. (Withdrawn) A composition comprising a cell wherein the cell comprises the vector of claim 39.
- 46. (Withdrawn) The composition of claim 47, wherein the cell comprises a neuron, glia cell, fibroblast, chondrocyte, osteocyte, endothelial cell, or hepatocyte.
- 47. (Withdrawn) The composition of claims 6, wherein the composition is in pharmaceutically acceptable form.
- 48. (Withdrawn) The composition of claims 6, wherein the composition is in an effective dosage.
- 49. (Withdrawn) The composition of claim 48, wherein the effective dosage is determined as a dosage that reduces the effects of Tay Sachs or Sandoff's disease.
- 50. (Withdrawn) A composition comprising an animal wherein the animal comprises the vector of claim 39.
- 51. (Withdrawn) A composition comprising an animal wherein the animal comprises the nucleic acid of claim 6.
- 52. (Withdrawn) A composition comprising an animal wherein the animal comprises the cell of claim 45.
  - 53. (Withdrawn) The composition of claim 50, wherein the animal is mammal.
- 54. (Withdrawn) The composition of claim 53, wherein the mammal is a murine, ungulate, or non-human primate.
- 55. (Withdrawn) The method of claim 54, wherein the mammal is a mouse, rat, rabbit, cow, sheep, or pig.
  - 56. (Withdrawn) The composition of claim 54, wherein the mammal is mouse.
- 57. (Withdrawn) The composition of claim 56, wherein the mouse comprises a HexB knockout.

- 58. (Withdrawn) The composition of claim 56, wherein the mouse comprises a HexA knockout.
- 59. (Withdrawn) The composition of claim 58, wherein the mouse further comprises a HexB knockout.
- 60. (Withdrawn) The composition of claim 54, wherein the mammal is a non-human primate.
- 61. (Withdrawn) A method of providing HEXA in a cell comprising transfecting the cell with the nucleic acids of claim 6.
- 62. (Withdrawn) A method of providing HEXB in a cell comprising transfecting the cell with the nucleic acids of claims 6.
- 63. (Withdrawn) A method of providing HEX- $\alpha$  and HEX- $\beta$  in a cell comprising transfecting the cell with the nucleic acid of claim 6.
- 64. (Withdrawn) The method of claim 63, wherein the step of transfecting occurs in vitro.
- 65. (Withdrawn) The method of claim 63, wherein the step of transfecting occurs in vivo.
- 66. (Withdrawn) A method of providing HEXS in a cell comprising transfecting the cell with the nucleic acids of claim 6.
- 67. (Withdrawn) A method of making a transgenic organism comprising administering the nucleic acid of claim 6.
- 68. (Withdrawn) A method of making a transgenic organism comprising administering the vector of claim 39.
- 69. (Withdrawn) A method of making a transgenic organism comprising administering the cell of claims 45.

- 70. (Withdrawn) A method of making a transgenic organism comprising transfecting a lentiviral vector to the organism at during a perinatal stage of the organism's development.
- 71. (Withdrawn) A method of treating a subject having Tay Sachs disease and/or Sandoff disease comprising administering the composition of claim 47.
- 72. (Original) A method of making a composition, the composition comprising a nucleic acid molecule, wherein the nucleic acid molecule is produced by the process comprising linking in an operative way a promoter element, an element comprising sequence encoding HEX-β, a IRES element, and an element encoding HEX-α.
- 73. (Original) The method of claim 72, wherein the HEX- $\beta$  element comprises a sequence having at least 80% SEQ ID NO:1 and the HEX- $\alpha$  element comprises a sequence having at least 80% to SEQ ID NO:3.
- 74. (Original) The method of claim 73, wherein any change in SEQ ID NO:1 or SEQ ID NO:3 is a conservative change.
- 75. (Original) The method of claim 72, wherein the sequence encoding HEX-β hybridizes to SEQ ID NO:2 under stringent conditions and wherein the sequence encoding the HEX-α hybridizes to SEQ ID NO:4 under stringent conditions.
- 76. (Withdrawn) A method of producing a composition, the composition comprising a cell, the method comprising administering the nucleic acid of claim 6 to the cell.
- 77. (Withdrawn) A method of producing a composition, the composition comprising a peptide, the method comprising expressing the nucleic acid of claim 6.
  - 78. (Withdrawn) The method of claim 77, further comprising isolating the peptide.
- 79. (Withdrawn) A method of producing a composition, the composition comprising an animal, the method comprising administering the nucleic acid of claim 6 to the animal.
  - 80. (Withdrawn) The method of claim 79, wherein the animal is a mammal.

- 81. (Withdrawn) The method of claim 80, wherein the mammal is a murine, ungulate, or non-human primate.
- 82. (Withdrawn) The method of claim 81, wherein the mammal is a mouse, rat, rabbit, cow, sheep, or pig.
- 83. (Original) A nucleic acid comprising a sequence encoding HEX- $\beta$  wherein the HEX- $\beta$  has the sequence set forth in SEQ ID NO:3, a sequence encoding HEX- $\alpha$ , wherein the HEX- $\alpha$  has the sequence set forth in SEQ ID NO:1, a promoter, and an IRES sequence, wherein the promoter is located 5' to the sequence encoding the HEX- $\beta$  and the sequence encoding the HEX- $\beta$  is orientated 5' to the IRES sequence and the IRES sequence is located 5' to the sequence encoding HEX- $\alpha$ .
- 84. (Original) A composition comprising a nucleic acid wherein the nucleic acid comprises a sequence encoding a first HEX-β and a sequence encoding a second HEX-β.
- 85. (Original) A composition comprising a nucleic acid wherein the nucleic acid comprises a sequence encoding a first HEX- $\alpha$  and a sequence encoding a second HEX- $\alpha$ .
- [86. (Original) A composition comprising four parts: 1) a promoter, 2) a sequence encoding a HEX- $\alpha$ , 3) a sequence encoding a HEX- $\beta$ , and 4) an integrated ribosomal entry site (IRES).
- 87. (Original) The composition of claim 6, wherein the promoter comprises a cell specific promoter.
- 88. (Original) The composition of claim 87, wherein the cell specific promoter comprises the Nuclear enolase specific (NSE) promoter.
- 89. (Original) The composition of claim 88, wherein the cell specific promoter comprises the sequence set forth in SEQ ID NO:69.

- 90. (Original) The composition of claim 87, wherein the cell specific promoter comprises the COLL1A1 promoter.
- 91. (Original) The composition of claim 90, wherein the cell specific promoter comprises the sequence set forth in SEQ ID NO:70 or SEQ ID NO:71.
- 92. (Withdrawn) A method of delivering a nucleic acid to a brain central nervous system cell comprising systemically administering a vector to the subject, wherein the vector transduces a blood cell, and wherein the blood cell fuses with a brain cell.
- 93. (Withdrawn) The method of claim 92, wherein the blood cell comprises a blood progenitor cell.
- 94. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for a blood progenitor cell.
- 95. (Withdrawn) The method of claim 92, wherein the blood cell comprises an endothelial cell.
- 96. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for an endothelial cell.
- 97. (Withdrawn) The method of claim 92, wherein the endothelial cell comprises a marker, wherein the marker is CD31.
- 98. (Withdrawn) The method of claim 92, wherein the blood cell comprises a microglia cell.
- 99. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for a microglia cell.
- 100. (Withdrawn) The method of claim 92, wherein the blood cell comprises a monocyte cell.

- 101. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for a monocyte cell.
- 102. (Withdrawn) The method of claim 92, wherein the blood cell comprises a macrophage.
- 103. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for a macrophage cell.
- 104. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker wherein the marker is CD11b.
- 105. (Withdrawn) The method of claim 92, wherein the blood cell comprises a lymphocyte cell.
- 106. (Withdrawn) The method of claim 92, wherein the blood cell comprises a marker for a lymphocyte cell.
- 107. (Withdrawn) The method of claim 105, wherein the lymphocyte cell comprises a marker wherein the marker is CD3.
- 108. (Withdrawn) The method of claim 92, wherein the brain cell comprises a purkinje cell.
- 109. (Withdrawn) The method of claim 92, wherein the brain cell comprises a marker for a purkinje cell.
- 110. (Withdrawn) The method of claim 109, wherein the marker is calbindin for Prkinje cerebellar cells
- 111. (Withdrawn) The method of claim 92, further comprising, adding the vector to a blood cell ex vivo producing a transduced blood cell, and administering the transduced blood cell to the subject.

- 112. (Withdrawn) The method of claim 111, wherein the blood cell comprises a blood cell obtained from the subject or is derived from a blood cell obtained from the subject.
- 113. (Withdrawn) The method of claim 111, wherein the blood cell comprises a progenitor cell.
- 114. (Withdrawn) The method of claim 111, wherein the blood cell comprises a marker for a blood progenitor cell.
- 115. (Withdrawn) A method of delivering a vector to a brain cell comprising, administering the vector to a subject, wherein the vector directly transduces the brain cell.
- 116. (Withdrawn) The method of claim 115, wherein the vector comprises the nucleic acid of claim 6.
  - 117. (Withdrawn) The method of claim 115, wherein the subject is a perinatal..
  - 118. (Withdrawn) The method of claim 115, wherein the subject is a neonatal.
- 119. (Withdrawn) The method of claim 115, wherein the brain cell is a brain cortex cell, a brain basal ganglia cell, a brain thalamus cell, a brain cerebellum cell, or a brain stem cell.
- 120. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises less than or equal to  $10^3$  infectious particles.
- 121. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises less than or equal to  $10^5$  infectious particles.
- 122. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises less than or equal to 10<sup>7</sup> infectious particles.
- 123. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises greater than or equal to 10<sup>3</sup> infectious particles.
- 124. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises greater than or equal to 10<sup>5</sup> infectious particles.

- 125. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises greater than or equal to 10<sup>7</sup> infectious particles.
- 126. (Withdrawn) The method of claim 115, wherein the administration of the vector comprises a m.o.i. of about 2.
- 127. (Withdrawn) The method of claim 116, wherein the vector reduces the inflammation of the brain.
- 128. (Withdrawn) The method of claim 116, wherein the vector reduces the deteriation of motor function due to a lysomal storage disease.
- 129. (Withdrawn) The method of claim 128, wherein the lysomal storage disease involves GM<sub>2</sub> gangliodisose.
  - 130. (Withdrawn) The method of claim 129, wherein the disease is Tay-Sachs disease.
  - 131. (Withdrawn) The method of claim 129, wherein the disease is Sandoff's disease.
- 132. (Withdrawn) A method of delivering a vector to a brain cell comprising systemically administering a vector to a perinatal subject.
- 133. (New) The composition of claim 7, wherein the HEX- $\beta$  has at least 70% identity to the sequence set forth in SEQ ID NO:3 and the HEX- $\alpha$  has at least 70% identity to the sequence set forth in SEQ ID NO:1.
- 134. (New) The composition of claim 133, wherein the integrated ribosomal entry site has the sequence set forth in SEQ ID NO:5.
- 135. (New) The composition of claim 7, wherein the HEX- $\beta$  has at least 85% identity to the sequence set forth in SEQ ID NO:3 and the HEX- $\alpha$  has at least 85% identity to the sequence set forth in SEQ ID NO:1.
- 136. (New) The composition of claim 135, wherein the integrated ribosomal entry site has the sequence set forth in SEQ ID NO:5.

- 137. (New) The composition of claim 136, wherein any change from SEQ ID NO:3 or SEQ ID NO:1 is a conservative change.
- 138. (New) The composition of claim 7, wherein the HEX- $\beta$  has at least 95% identity to the sequence set forth in SEQ ID NO:3 and the HEX- $\alpha$  has at least 95% identity to the sequence set forth in SEQ ID NO:1.
- 139. (New) The composition of claim138, wherein any change from SEQ ID NO:3 or SEQ ID NO:1 is a conservative change.
- 140. (New) The composition of claim 138, wherein the integrated ribosomal entry site has the sequence set forth in SEQ ID NO:5.
- 141. (New) The composition of claim 140, wherein any change from SEQ ID NO:3 or SEQ ID NO:1 is a conservative change.